# 1 Program 3drefit

## 1.1 Purpose

Changes some of the information inside a 3D dataset's header. Note that this program does NOT change the .BRIK file at all; the main purpose of 3drefit is to fix up errors made when using to3d. To see the current values stored in a .HEAD file, use the command '3dinfo dataset'. Using 3dinfo both before and after 3drefit is a good idea to make sure the changes have been made correctly!

## 1.2 Usage

3drefit [options] dataset ...

## 1.3 Options

#### -orient code

Sets the orientation of the 3D volume(s) in the .BRIK. The code must be 3 letters, one each from the pairs {R,L} {A,P} {I,S}. The first letter gives the orientation of the x-axis, the second the orientation of the y-axis, the third the z-axis:

 $egin{array}{lll} R &= {
m right-to-left} & L &= {
m left-to-right} \\ A &= {
m anterior-to-posterior} & P &= {
m posterior-to-anterior} \\ I &= {
m inferior-to-superior} & S &= {
m superior-to-inferior} \\ \end{array}$ 

\*\* WARNING: when changing the orientation, you must be sure to check the origins as well, to make sure that the volume is positioned correctly in space.

- -xorigin distx
- -yorigin disty
- -zorigin distz

Puts the center of the edge voxel off at the given distance, for the given axis (x,y,z); distances in mm. (x=first axis, y=second axis, z=third axis). Usually, only -zorigin makes sense. Note that this distance is in the direction given by the corresponding letter in the -orient code. For example, '-orient RAI' would mean that '-zorigin 30' sets the center of the first slice at 30 mm Inferior. See the to3d manual for more explanations of axes origins.

\*\* SPECIAL CASE: you can use the string 'cen' in place of a distance to force that axis to be re-centered.

- -xdel dimx
- -ydel dimy
- -zdel dimz

Makes the size of the voxel the given dimension, for the given axis (x,y,z); dimensions in mm.

\*\* WARNING: if you change a voxel dimension, you will probably have to change the origin as well.

#### -TR time

Changes the TR time to a new value (see 'to3d -help').

\*\* WARNING: this only applies to 3D+time datasets.

#### -newid

Changes the ID code of this dataset as well.

#### -nowarp

Removes all warping information from dataset.

#### -statpar v ...

Changes the statistical parameters stored in this dataset. See 'to3d -help' for more details.

### -markers

Adds an empty set of AC-PC markers to the dataset, if it can handle them (is anatomical, is in the +orig view, and isn't 3D+time).

\*\* WARNING: this will erase any markers that already exist!

#### -view code

Changes the 'view' to be 'code', where the string 'code' is one of 'orig', 'acpc', or 'tlrc'.

\*\* WARNING: The program will also change the .HEAD and .BRIK filenames to match.

If the dataset filenames already exist in the '+code' view, then this option will fail. You will have to rename the dataset files before trying to use '-view'. If you copy the files and then use '-view', don't forget to use '-newid' as well!

### -byteorder bbb

Sets the byte order string in the header. Allowable values for 'bbb' are:

LSB\_FIRST MSB\_FIRST NATIVE\_ORDER

Note that this does not change the .BRIK file! This is done by programs 2swap and 4swap.

#### -appkey ll

Appends the string 'll' to the keyword list for the whole dataset.

### -repkey ll

Replaces the keyword list for the dataset with the string 'll'.

#### -empkey

Destroys the keyword list for the dataset.

#### -type

Changes the type of data that is declared for this dataset, where 'type' is chosen from the following:

#### ANATOMICAL TYPES

```
== Spoiled GRASS
                            fse
                                      Fast Spin Echo
                                 ==
spgr
      == Echo Planar
                                      MRI Anatomy
                            anat
epan
                                 ==
ct
      == CT Scan
                                 == SPECT Anatomy
                            spct
pet
      == PET Anatomy
                            mra
                                 ==
                                      MR Angiography
      == B-field Map
                            diff
                                      Diffusion Map
bmap
      == Other MRI
                                      Anat Bucket
omri
                            abuc ==
```

#### FUNCTIONAL TYPES

$_{ m fim}$	==	Intensity	$\operatorname{fith}$	==	Inten+Thr
fico	==	Inten+Cor	$_{ m fitt}$	==	Inten+Ttest
fift	==	Inten+Ftest	fizt	==	Inten+Ztest
fict	==	Inten+ChiSq	$\operatorname{fibt}$	==	Inten+Beta
fibn	==	Inten+Binom	$\operatorname{figt}$	==	Inten+Gamma
fipt	==	Inten+Poisson	fbuc	==	Func-Bucket

The options below allow you to attach auxiliary data to sub-bricks in the dataset. Each option may be used more than once so that multiple sub-bricks can be modified in a single run of 3drefit.

```
    -sublabel n ll Attach to sub-brick #n the label string 'll'.
    -subappkey n ll Add to sub-brick #n the keyword string 'll'.
    -subrepkey n ll Replace sub-brick #n's keyword string with 'll'.
    -subempkey n Empty out sub-brick #n' keyword string
```

### -substatpar n type v ...

Attach to sub-brick #n the statistical type and the auxiliary parameters given by values 'v ...', where 'type' is one of the following:

$_{\mathrm{type}}$	Description	PARAMETERS
fico	Cor	Samples, Fit-Parameters, Ort-Parameters
fitt	Ttest	Degrees-of-Freedom
fift	Ftest	Numerator and Denominator Degrees-of-Freedom
fizt	Ztest	N/A
$\operatorname{fict}$	ChiSq	Degrees-of-Freedom
fibt	$\operatorname{Beta}$	Alpha and Beta (exponenents)
$\operatorname{fibn}$	Binom	Number-of-Trials and Probability-per-Trial
$\operatorname{figt}$	Gamma	Shape and Scale
$\operatorname{fipt}$	Poisson	Mean